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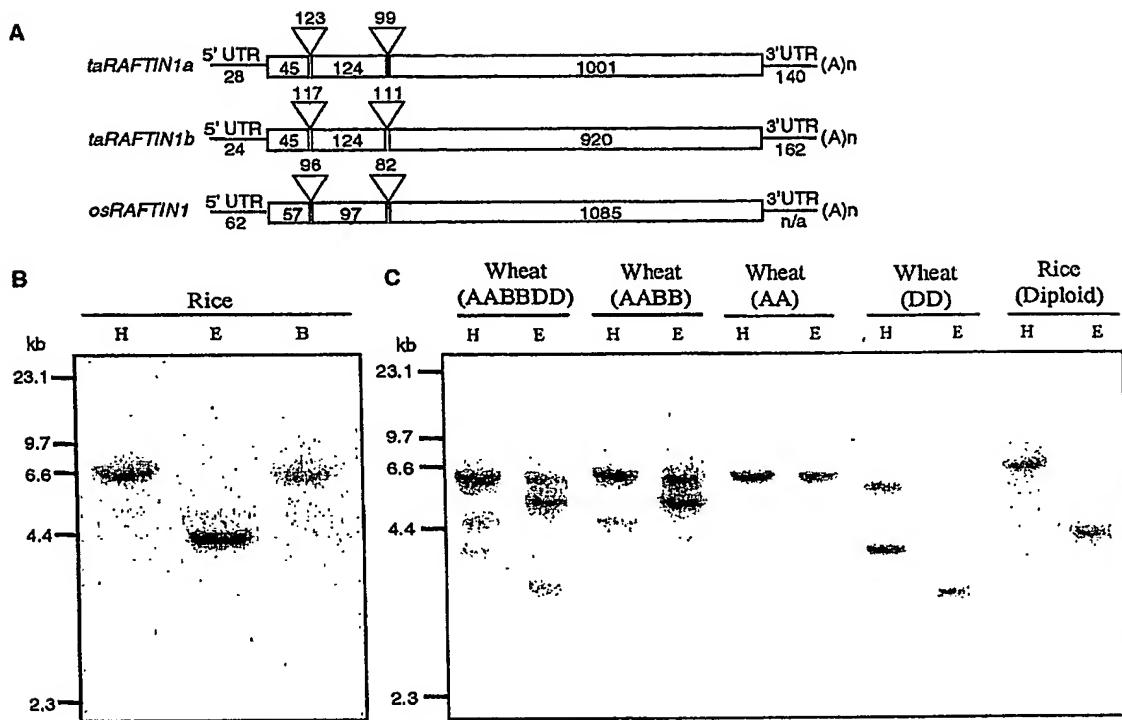
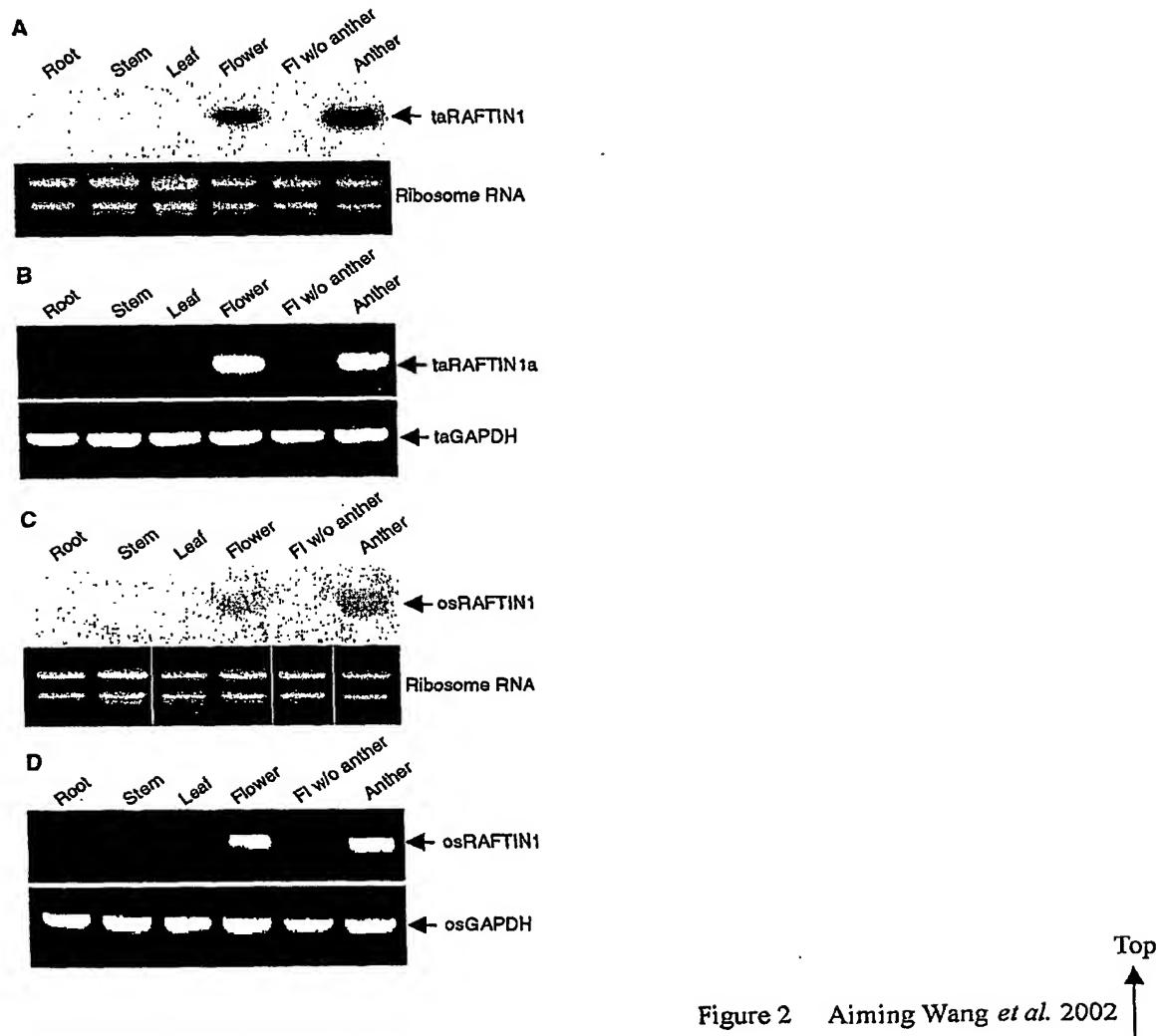


Figure 1

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Figure 2 Aiming Wang *et al.* 2002

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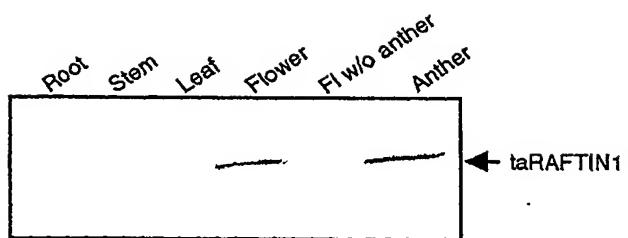


Figure 3 Aiming Wang *et al.* 2002

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A

taRAFTIN1a 1 MARELVALLATTLVAVOAGGQLGHAAPATAEVEWRALPHSPLPAVLRLLKQPAAGVELLTEATSFVRDAEDR~~RA~~PED 78
 taRAFTIN1b 1 ~~RA~~-----G-----X-----S-----P-----X----- 72
 osRAFTIN1 1 ~~RA~~-----LVAVA-AA-VLS-D-----S-----F-----RPDTSF-VGKA-A~~RA~~AGAARTGF--- 74

taRAFTIN1a 79 YRDYSRSPPDDEPSKSTGAA~~AA~~SGARD~~FDY~~DDYS~~AA~~GGDKLRGAASGAR~~AAA~~DFDYDDYS~~GA~~DKLRGAT~~AAA~~ 141
 taRAFTIN1b 73 -----SS-----V-~~RA~~-----~~AAA~~-----~~AAA~~-----~~AAA~~-----N-----ER-----~~AAA~~ 114
 osRAFTIN1 75 T--RG-DSPTTA-GLDL-GDFGE~~PAP~~-G-A-AQGE-GGGAA-A-EQVLAVDAG-N-K-V-R-L-GSSTAGGE 153

taRAFTIN1a 142 ~~AAA~~EEYKAPSSLAGNGASMARG~~G~~KAET~~TT~~V~~EE~~AVRGKRLP~~FP~~PPAT~~PA~~LGFL~~PRQV~~ADSV~~PF~~TTAALP 212
 taRAFTIN1b 115 ~~AAA~~-----S-Y-----X-----R-----H----- 185
 osRAFTIN1 154 NDDEPFGYD-----GSGTAA~~TT~~V-TGA-----E-----Y-A-TS-----R-----I-----A----- 233

taRAFTIN1a 213 ~~GV~~LATE~~GV~~ASDSATVASMEATL~~R~~ACESPTIAGESKFCATS~~LE~~ALVERAMEVLGTRDIRPVT~~STL~~PRAGAPLQTYTVRSVR 292
 taRAFTIN1b 186 ~~I~~-----T-P-----G-----VA-Q 265
 osRAFTIN1 234 ~~A~~-----L-----P-T-EA-G-RE-----T-W-L-----G-A-----AALA-----G-----A-A-A-L 313

taRAFTIN1a 293 PVEGGPVFVACHDEA~~Y~~PTVYRCHT~~T~~GPSRAYMVDMEG~~AA~~ARGGDAV~~T~~IA~~T~~VCH~~T~~DT~~S~~LWN~~P~~HVS~~E~~KLLG~~T~~KPGG~~P~~V 369
 taRAFTIN1b 266 -----T-----AAA-----A-----A----- 342
 osRAFTIN1 314 -----AG-----Q-----A-----E-----DGGGD-----E-----V-----N-R-----S----- 392

taRAFTIN1a 370 CHLMPYGHIIWAKNVNRSPA 389
 taRAFTIN1b 343 -----K----- 362
 osRAFTIN1 393 -----V-----KS-T----- 412

B

taRAFTIN1a 169 ~~FF~~HEEAVRVGKRLP~~FR~~FP~~PA~~T~~AA~~ALGFL~~PRQV~~ADSV~~PF~~TTAALPGV~~LA~~TE~~GV~~ASDSATVASMEATL~~R~~ACESPTIAGES 246
 PG-bet 415 --R-KMLKS-TIMPM~~A~~-DIKDKMPKRS---VI-SKL---S-SKIAELKAI~~F~~HAGDE-Q~~VE~~KMIGDA-SE--RAPS---T 492
 RD22 176 --L-KDLVR--EMNV--NAEDG~~Y~~GGKTA---GE-ET---GSEKFSET-KR-S-EAG-EEAEM-KK-IEE--ARKVSG-E 255
 ASG-1 87 --N-HD-LE--TE-MY--SV~~A~~-K---Q-R-VQ~~E~~I---S-R-ADI--L-HIPPG-SEAADVAT--GL-DAAAHGDVV 164
 CFC1 124 --L-KDMHP-ATMSLH-TEN-~~AA~~AKSA--Y-T-QK~~I~~--SSDK--E~~IF~~NK-S-KPG-LK-EM-KN-IKE--Q-A-E--E 200
 SCB1 92 --L--DL-A--IFNMK-VNN-KA-TVPL---ISK~~Q~~I--SEDKKKQ--ML--EAN-SNAKIIAE--GL-QE-ATEG-R 169

taRAFTIN1a 247 KFCATS~~LE~~ALVERAMEVLGTRDIRPVT~~STL~~PRAGAPLQTYT~~A~~RSVR~~P~~VEGGPV~~V~~ACHDEA~~Y~~PTVYRCHT~~T~~GPSRAY 324
 PG-bet 493 --R-VN-A-DMIDFATS--RN~~V~~~~A~~-RT-EDTKGSNGNIMIGSVKG~~ING~~GGKVTK~~S~~-S-CTL---LL-Y--SV~~PKV~~-V- 569
 RD22 256 --Y-----SM-DFVSK--KYHV-A-STEVAKKN--M-K-XIAAG-KKLSDDK~~S~~-V--KQK--FA-FY--KAMMTV- 334
 ASG-1 165 R~~A~~-V--PDDM-G--AA---SNMQVLAPS--TG-MS--P--~~AA~~-A-K--D-SQ~~A~~--G--P-L---S-----SVQTGT- 241
 CFC1 201 --Y-----SMIDYSISK--KV-~~AA~~Q-STEVEQ~~A~~TPM-K--IAAAG-QKMTDDK~~A~~-V--KQ~~N~~-A-A-FY--KSETT-- 276
 SCB1 170 --H-----SM-DFV~~V~~SA--K~~VG~~AFSTEKERETES~~A~~GKFV-VKNG--KLGDDHV~~A~~IA--PMS---V-FG--LVPR-SG- 246

taRAFTIN1a 325 MVDMEGARG~~G~~DAV~~T~~IA~~T~~VCH~~T~~DT~~S~~LWN~~P~~HVS~~E~~KLLG~~T~~KPGG~~P~~CHLMPYGHII 380
 PG-bet 570 EA-ILDPNSKV~~K~~INHG~~V~~AI--V---S-G-S-GA-VA--SG--KIE---WIFENDMTW 626
 RD22 338 A-PL--~~AA~~EN-MRAKAVA--KN--A--N-LA--V-KV--TV---FL-ET-VV- 389
 ASG-1 242 VME-QSSY-N-G-LKLVA--RN-TS-D-----V-AS---L-I--FV---V-F 298
 CFC1 277 --PL--~~AA~~AD-TKA~~K~~AVA-----A--K-LA-QV-KV~~E~~--TI---FL-RD--V- 331
 SCB1 247 L-RLK-~~AA~~E~~D~~-VR-KAVVA--R--K-DHN-GA--V-NL---NGT---V~~F~~E-NLL- 301

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Figure 4 Aiming Wang *et al.* 2002

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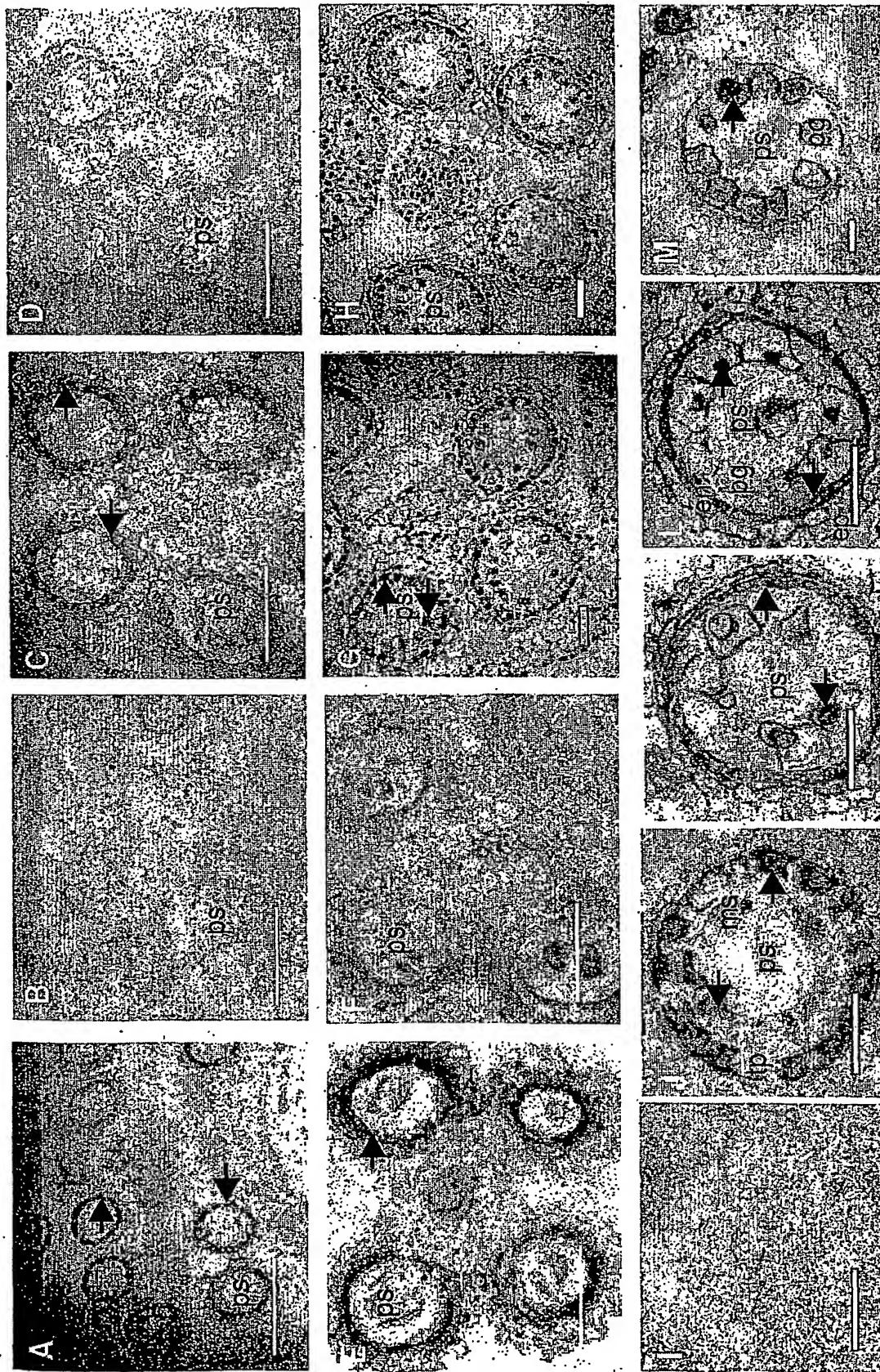


Figure 5

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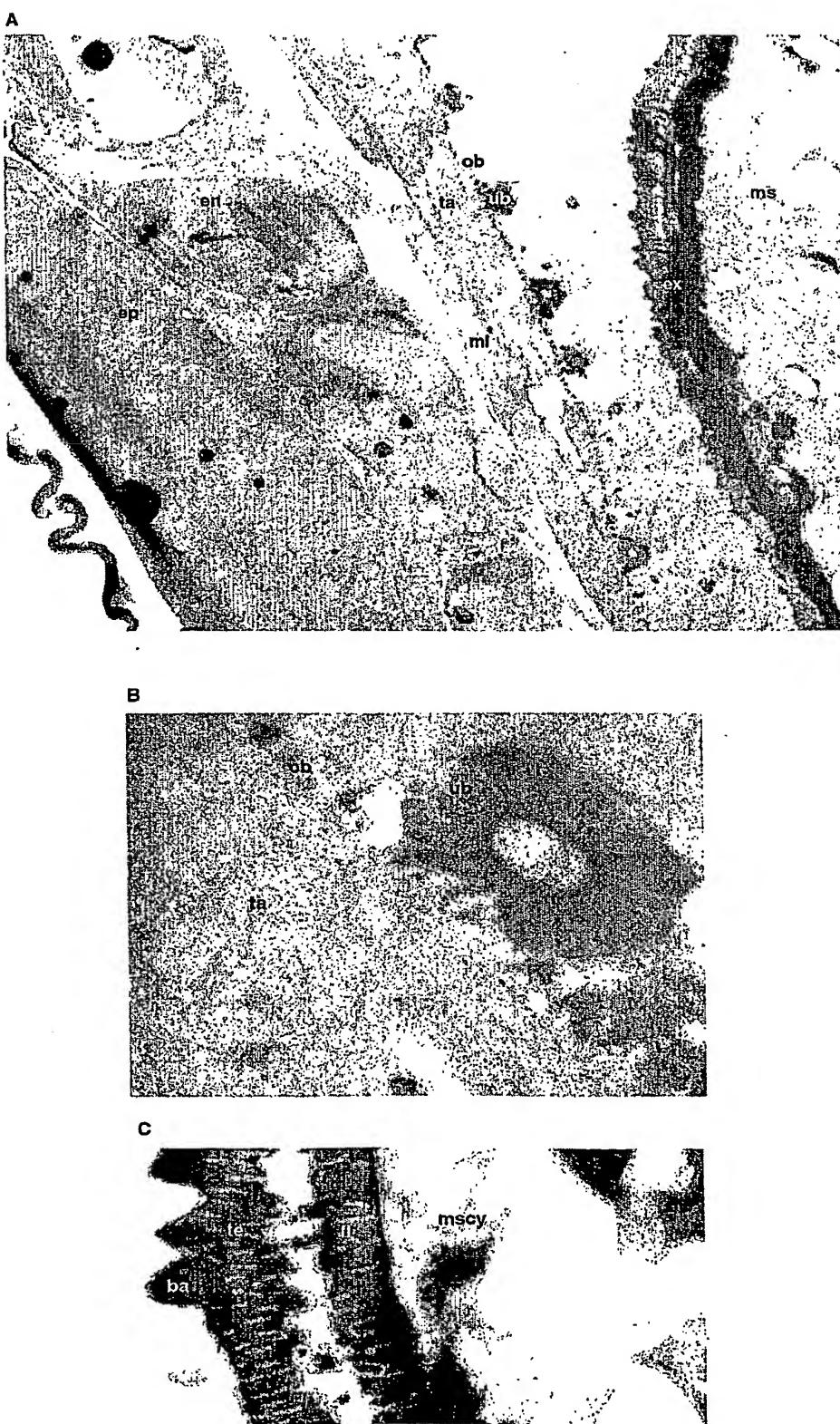
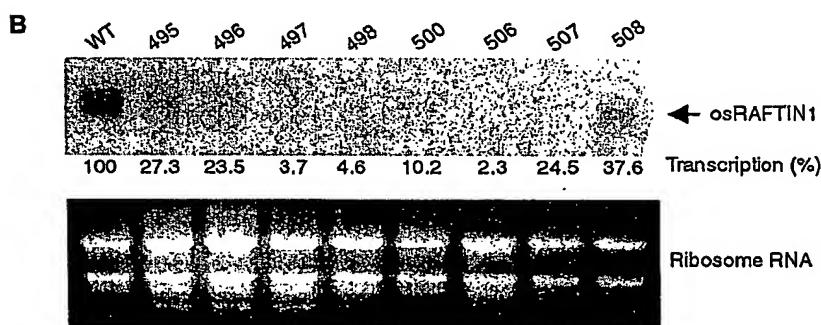
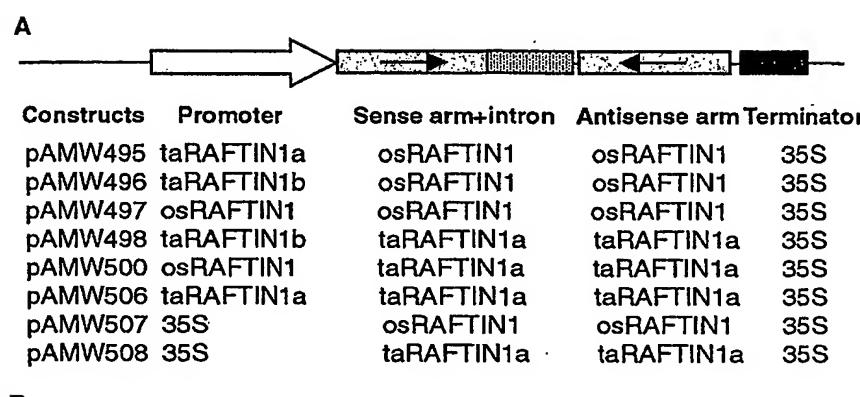
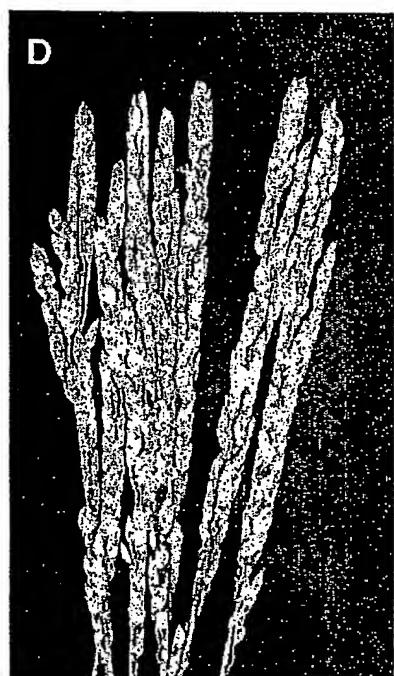
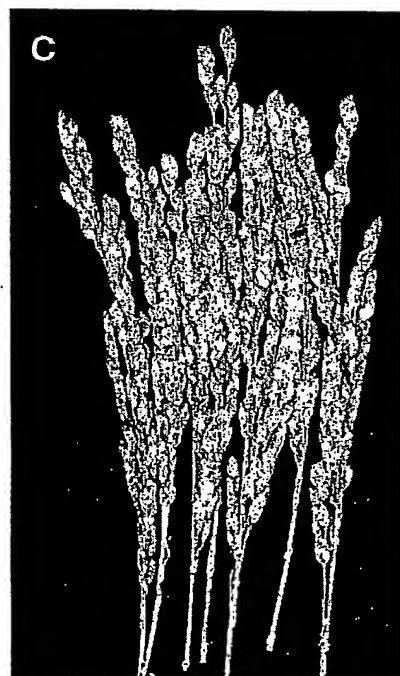
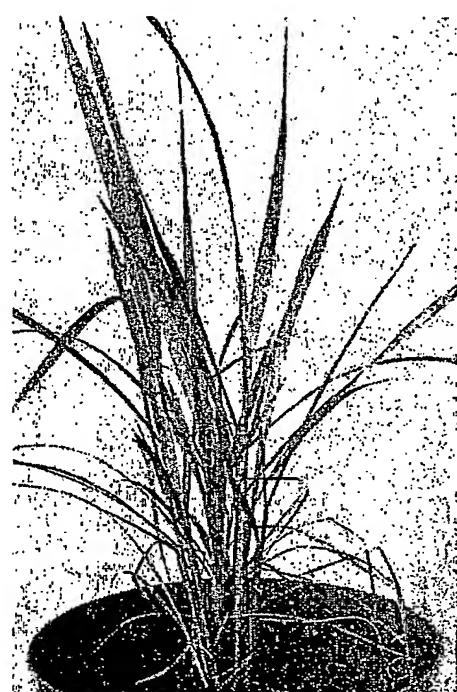


Figure 6

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Figure 7 Aiming Wang *et al.* 2002

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Figure 8ABCD Aiming Wang *et al.* 2002

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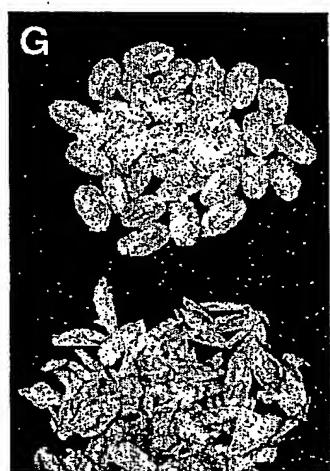
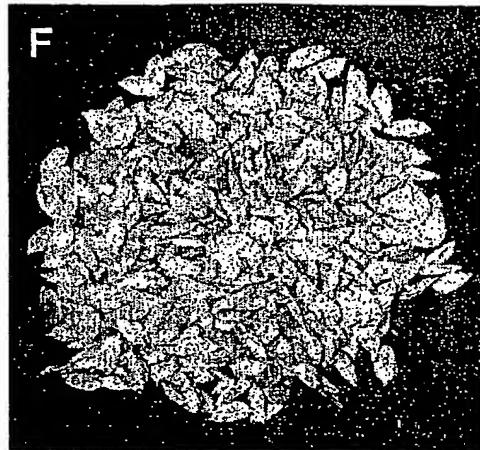
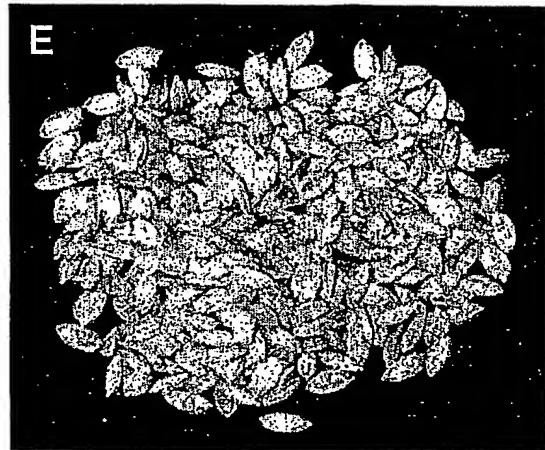


Figure 8EFGH Aiming Wang *et al.* 2002

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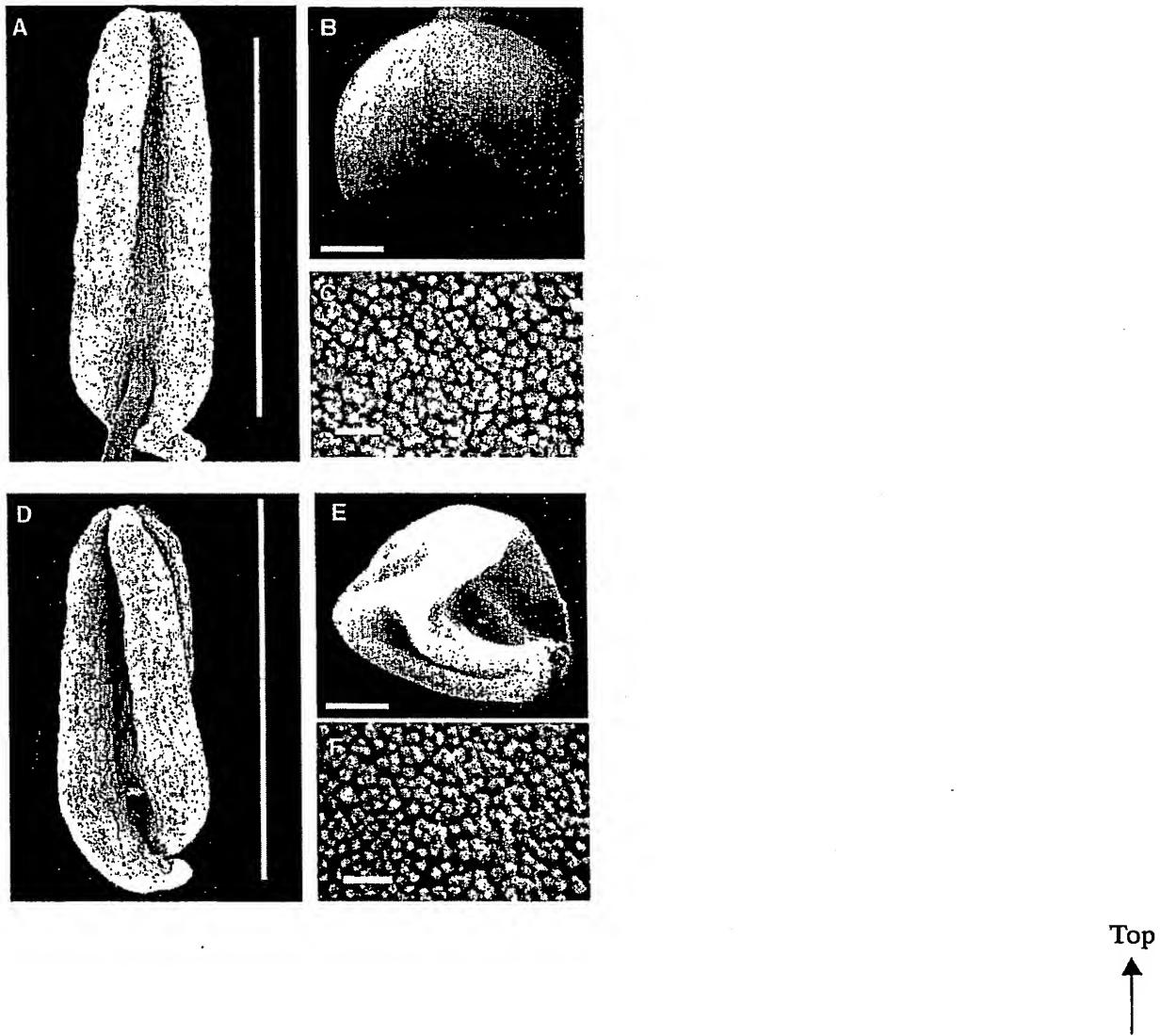
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Figure 8I Aiming Wang *et al.* 2002

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Figure 9 Aiming Wang *et al.* 2002

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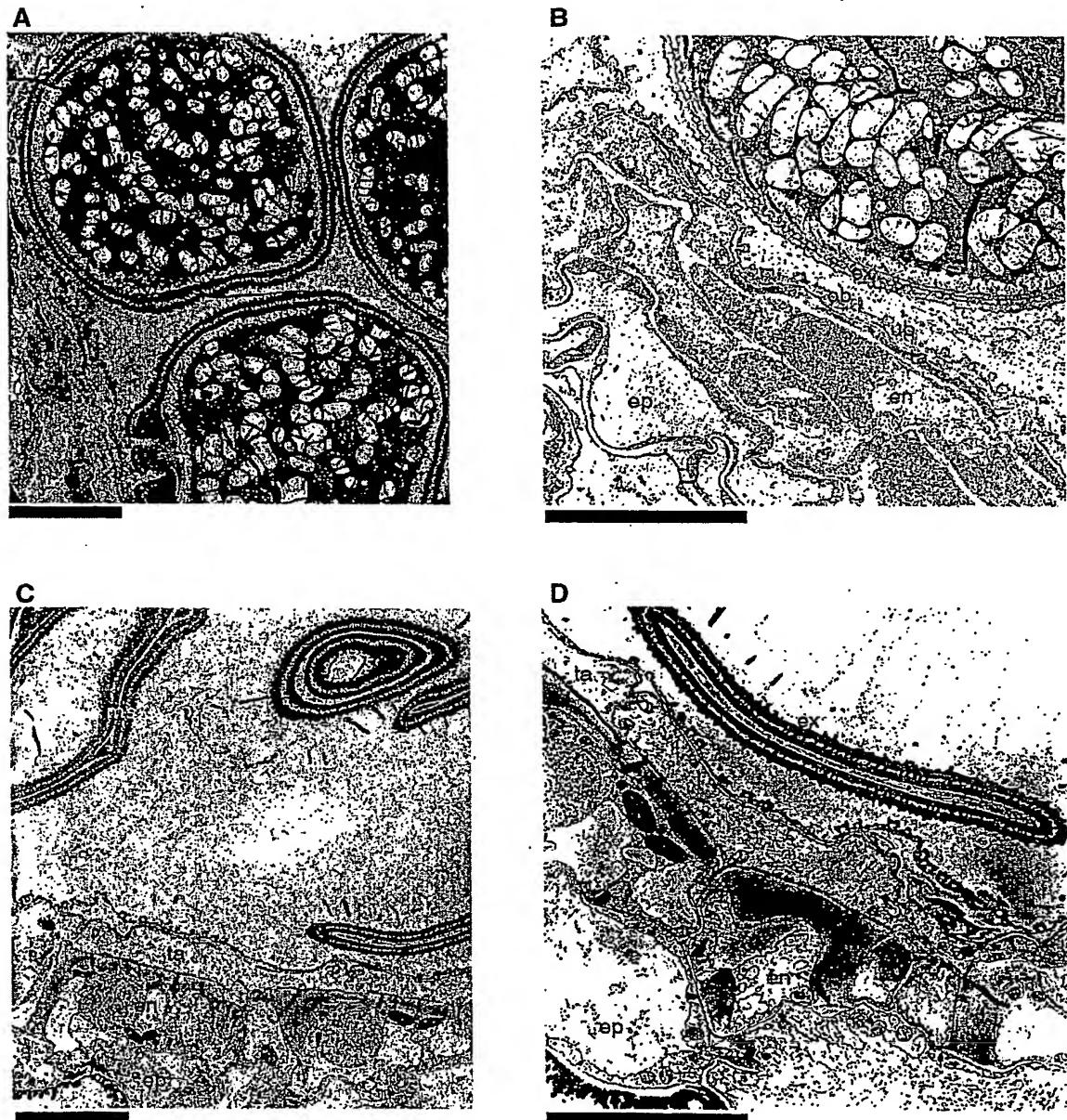


Figure 10

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Fig. 11. *taRAFTIN1a* cDNA sequence (1338 nt excluding the polyA tail, ORF from nt 29 to nt 1198). Start codon and stop codon are underlined.

CTCTGGACCTCTCACCTAGCGCACATCCATGGCGCGCTCCTCGCCCTCCTCGCCACCAC
CCTGGTCGCGGTTCAGGCTGGAGGGCAGCTGGGCCACGCGGCCGGCGACGGCGGAGGTGTT
CTGGCGCGCCGTGCTGCCACACTCGCCATTGCCGACGCCGTTCTCCGCCTCTCAAACAAACC
CGCAGCAGGTGTTGAACTGCTCACAGAACGCCACCAGCTCGTGAGGGATGCGGAGGACAGGCC
CCCCTCGACTACCGTGATTACAGCCGCTGCCGCCGATGATGAACCGAGCAAGAGCACCGG
CGCCGCCTCCGGGGCGCGGGACTTCGACTACGACACTACAGCGGGGGCGACAAGCTCCGTGG
CGCCGCCTCCGGGGCGCGGGACTTCGACTACGACACTACAGCGGGGGCGACAAGCTCCGTGG
CGCCACCGATGAATACAAGGCCCGAGCAGCAGCCTCGCTGGAAACGGGGCGTCCATGGCTAG
GGGCGGCAAGGCGGAGACGACGACGGTGTTTCACGAGGAGGCGGTGCGGTCGCAAGAG
GCTCCCATTCGCTCCGCCGGCGACTCCCGCCGCGCTCGGTTCCTCGCGCCAGGTCGC
CGACTCCGCTCCGTTTCACGACGGCCCGCTGGCGTCCTCGCGACGTTCGGCGTCGCGT
CGACTCCGCCACGGTGGCCAGCATGGAGGCGACGCTCGCGCCTCGGAGTCGCCACATCGC
CGGGGAGTCCAAGTTCTGCGCGACCTCGCTGGAGGCCCTGGTGGAGCGCGCCATGGAAGTGCT
GGGGACCCCGGACTAGGCCGGTGACGTCGACGCTGGCCCCGGGCGCCGGCTGCAGAC
GTACACCGTCGCTCCGTCCGTGGGGCCGGGGTGAGGGGGGCCGTCTTCGGGCGTGCCCACCGACGA
GGCCTACCCGTACACCGGTACCGGGTCCACCCACACTGGCCCGTCCCAGGGCGGTACATGGGTGGA
CATGGAGGGCGCGCGGGCGACGGCGTGACCCATCGCCACCGGTGTGCCAACACCGACACGTC
CCTGTGGAAACCCGGGAGCACGTCTCCTCCAAGCTCCGGACCAAGCCTGGCGGCACGGGT
CTGCCACCTCATCGCGTACGGGCACATAAATCTGGGCCAGAACGTGAATCGCTCGCCGGCG
GTAGCGGGCCGGGCAGCTGTGGTCTCGCCGGAACTAAAGATCGATGTACTACTACTACTATCTG
TTTCCACCTACGTCTCTGTGGTTCAGACCACCAGATGGTCACCAGAGCAGCGCGTTGTAAATAA
AAGAACAGCTTCTGCAAAAAAAAAAAAAA

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Fig. 12. *taRAFTIN1a* genomic sequence (1560 bps including two introns). Introns are shown in lower case letters. Start codon and stop codon are underlined.

CTCTGGACCTCTCACCTAGCGCACATCCATGGCGCGCTTCCTCGCCCTCGCCACCAC
CCTGGTCGCGtaatggccgaagaaggcactgagcaacgcctgcatcttcttcatttcggaa
actgcaccttagtgcatttcgcatgagattgatcgatcacaaactggtgctaacggcctgtttc
gtcacagGTTCAGGCTGGAGGGCAGCTGGGCCACGCGGCCGGCAGCGCGGAGGTGTTCTG
GCGGCCGTGCTGCCACACTCGCCATTGGCCGACGCCGTTCTCCGCCTTCTCAAACAAACCGC
AGCAGgtctgtcttcatgttccttcctgtcgccctccgttaactgtcttcttctcgag
tttgattgaccgcaaacacaaaaaaatgcatgcacgcacagGTGTTGAACTGCTCACAGAAGC
CACCAGCTCGTGAGGGATGCCGAGGACAGGCCCCCTCGACTACCGTGATTACAGCCGCTC
GCCGCCGATGATGAACCGAGCAAGAGCACCGCGCCCTCCGGGGCGCGGGACTTCGACTA
CGACGACTACAGCGGGGGCGACAAGCTCCGTGGCGCCCTCCGGGGCGCGGGACTTCGACTA
CGACGACTACAGCGGGGGCGACAAGCTCCGTGGCGCCACCGATGAATACAAGGCCCGAGCAG
CAGCCTCGCTGGAAACGGGGCGTCCATGGCTAGGGCGGCAAGGGCGGAGACGACGACGGTTT
CTTCACGAGGAGGCGGTGCCGGTCGCAAGAGGCTCCATTCCGCTTCCGCCGGCGACTCC
CGCCCGCTCGTTCCCTGCCGCCAGGTCGCCACTCCGTCCGTCCCGACT
GCCTGGCGCTCTCGCGACGTTCGCGGTCGCGACTCCGCCACGGTGGCCAGCGATGGAGGC
GACGCTCGCGCCCTCGGAGTCGCCACCGCCGGGGAGGTCCAGTTCTCGCGACCTCGCT
GGAGGCCCTGGAGCGCGCCATGGAAGTCGGTGGGGACCCCGACATCAGGCCGTGAGTCC
GACGCTGCCCCGCCGGCGCCCCCGCTCGACAGGTACACCGGTCCGTCGCCGGGTGGGA
GGGGGGCCCTGTCTCGGTGGCGGTCCACCGACGGGCCTACCGGTACACCGGTACCCGGGTGCCA
CACCACTGGCCCGTCAGGGCGGTACATGGGTGGACATGGGAGGGGCGCGCCGGCGACGGCGGT
GACCATCGCCCACCGGTGTGCCACCGACACGACGTCCCTGTGGAACCGGAGCAGTCTCCTTCAA
GCTCCCTGGGCACCAAGCTGGCGGCACGCCCGGGTCTGCCCACCTCAGCCGGTACGGGCACATAAT
CTGGGCCAAGACGTGAATCGCTGCCCGGCGGTGAGCGGCCGGGCAGCTGTGGGTCTCGCCG
GAACTAAAGATCGATGTACTACTACTACTATCTGTTTCCACCTACGTCTGTGGGTCTCGCCG
ACCAGATGGTCACCACCAGAGCGACCGGTTAAAAGAAACAGCTCTGC

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Fig.13. *taRAFTIN1a* promoter sequence (1719 bps).

CTGTCGATGGCGCTCTGCTTGTGATTCTTCTTAGGGAACCTCGTCTCTGGGCCTCCGAGG
CCTGCAACCCTGTATCAGGACAATTCTGACTGGCCTCCAGGAGTCCTAACAGCCACCGACCTG
GTCCACTGGGCCATCTAGAGTATCTTGAAGTGTGTTGCACAAATCCCGCTAATTAAGGGA
TGTGATGATGATGGTTCTGAATCCGCGCCTTACCTCGAAAACGGGAAATTGCAAAGGAT
ATATGGCACCTGTCGCGTGTGAGGCCAGACGCTTCGGTTCAAGCTGGTTATAGGGAGGGGG
AAACGAAGGGTTTTCTCCCTCTGTCTTCATCCATTTCGTCTCCCAGCCCTAGCTCCCAA
AAGCGTGTGCCACCTCAAAGTCTTCAGCGCTGCTCACGTAGCCCCCGTCCACCCCTCCT
GCCACCAAGATGGCCCGAACCAAGAGCGAGAAGGTTCTAAAGGTTCCCAGCTAGGATCTGCC
GCCGCTGGAACGGGCTGAAGCGGAAGAGGGTCGCTCCAAGGGTGGTATGAAACAACAGCCG
GAAGCCCCAAGACTACAGGAAAGTGGTTCCCTCCTCGGCCACCGACAAAAAAACTTCAGGGT
CTCGTGGAGATAGGGCTGATGCCAGGGATTGGAGTGCCGCTCCGGGGACGAGGCTCCG
CCAACTCCTCGCGACGGTGAGCACATCCTCTGCCCTGGAGTATAATTTCGGAGGGGCTCGGG
TTTCCCCTACACGACTTCGTTGCCGGATCTTGCCTTACGGCTGCTAGCTACACCACATC
CCGTCAAACGGGTTCTTACATTGCAAACCTTCATCACATTGCGAGTGCTTCTCGGGACT
GCCGCTCACTTAAGTTGTTCAATACTTCAATCAGGACTGCGTTCAGACCAACGGGACATC
GTCTACGACCCCGAACACCAAATTCTCGCCACATACCTCCGAAAATAATCCTATAACACC
TGGTCTCACGCTTACATCTCGTAAGATTGCCATGTGTACTTCACCAATCTGATGCATCCCTT
TTTCCCCAAGATTATATGCCTGATCTGTATTGCTCCGCTTTCGAGATTGATGTTA
ATTGATGAAGCCAAGCAATCCGGATGCCGTCGGTGCACTAGATGGCTAGCTTCTACGG
TGCTGGGCCTGCCGGCGAGGGGCGCAGGCCACGTAGGAGACTGTTAGGATTCTGGGCTGG
ACCGTGGTGGCGTGAAGTTGGGAAGGAGGATTGAGGAAGAAGGATGCATCAAGATTGGTGA
GAACACGTGGCATCCTCTAGAGTAGGTCTTACGAGATGAAGCCTGAGACCAGGTCGTATGGGA
TTATTTCCGGACCTCCCGAACGCCGAAAGCTTAACTGCAGCTGCGTGGACGGCGAGCACC
GCACCGCACACGAACCGAACCTGACGCTGCCGCCACACAAACAGCCATTGCGCGCGGAT
CGTCGGATGTACGCCAGGATTATATTCTCCGGTGCCGACGTACCATGCGATCGCACAGCT
CACGTGAGAGCTTCTGTTGGCGTCGCCGTCAATGAAACACCTCCGTCGAGCCGACGA
CGCCTATAAGTACCTCGTCTGATCGCATCATCACTCCCAAGTACTACAACCTCTGGACCTCTC
ACCTAGCGCACATCCCATG

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Fig.14. *taRAFTIN1b* cDNA sequence (1275 bps excluding the polyA tail, ORF from nt 25 to nt 1113). Start codon and stop codon are underlined.

CGACCTCTCACCTAGCGCACATCCATGGCGCGCTTCCTCGTCGCCCTCCCTCGCTGCCACCCCTG
GTCGCGGTTCAAGGCTGGAGGGCAGCTGGGCCACGCGGCCGGCTACGGGGGAGGTGTTCTGG
CGCGCCGTGCTGCCGCACTCGCCATTGCCTGACGCCGTTCTCCGCCTCCCAAACAACCTGCA
GCAGAAATCCACCAGCTCGTAGAGAGACCCGAGGACAGGCCCCCTCGACTACCGTGATTAC
AGCCGCTCGTCCGATGATGAACCGAGCAAGAGCACCGTCGCCGCTCCGGAGCGGGGGC
TTCGACTACGACAACACTACAGCGGGGCGACGAACGTCGTGGTGCCACCGATGAATACAAGGC
CCGAGCAGCAGCCTCGCTGGAAGCGGGCGTACATGGCTAGGGCGGCAAGGCAGACGACG
ACGGTGTCTTCACGAGGAGGCGGTGCGCTCGGCAGGAGGCTCCATTCCACTTCCCAGCG
GCGACTCCCGCGCTCTCGGTTCTGCCGCGCAGGTCGCCGACTCCGTCCGTTACGACG
GCCGCCTGCCGGCATCCTCGCAGCTTGGCATCGCTCCGACTCCACACGGTGCCAGC
ATGGAGGGGAGCAGCTGCGCGCTGCGAGTCGCCACCATGCCGGGAGTCCAAGTTCTGCGCG
ACTTCGCTGGAGGCCCTGGGGAGCGCGCCATGGGAGTGCTGGGGACCCGGGACATCAGGCCG
GTGACGTCGACGCTGCCCGCGCCGGCGCCCGCTGCGACAGTACACCGTCGCGCGTGCAG
CCGGTGGAGGGGGGGCCTGTTCTCGTGGCGTGCCACGACGAGGCCTACCCGTACACCGTGTAC
CGGTGCCACACCACCGGCCGTCCAGGGCGTACACGGTGGACATGGAGGGCGCGCGGCC
GACCGGGTACCATGCCGCCGTGCGCACACCGACACGTCCCTGTGGAACCCGGAGCACGTC
TCCTCAAGCTCCTCGGCACCAAGCCGGCGCACGCCGGTCTGCCACCTCATGCCGTACGGG
CACATAATCTGGGCCAAGAACGTGAAGCGCTGCCGGCGTGAACGGCCTTGCAGCTCTGTT
GTCGCCGGAACTAAGATCGATGTACTACTACTATCTGTTCTACCTACGTCTTCTGTTGTT
ATACCAACAGATGGTCACCCAAAGAGCAAGCGTTGTAATAAAAGAACAGCTTTTGAGAAG
CTGGTGTCTTATTTAAAAAAAAAA

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Fig.15. *taRAFTIN1b* genomic sequence (1503 bps including two introns). Introns are shown in lower case letters. Start codon and stop codon are underlined.

CGACCTCTCACCTAGCGCACATCCATGGCGCGCTTCCTCGTCGCCCTCGCTGCCACCTG
GTCGCGgtaatggccgaagaagagcaacgcctgcatcttcttcatttggcaaattgcaccta
gtacattttgcataatcaatcacaactggtgctaacggcctgttgcgtccag
GTTCAAGGCTGGAGGGCAGCTGGGCCACGCCGGCGCTACGGGGGAGGTGTTCTGGCGCGCC
GTGCTGCCGCACTGCCATTGCCTGACGCCGTTCTCCGCCCTCAAACAAACCTGCAGCAGgt
ctgtcttgcataatcctcgtcgccctccgttaactgtcttcttctcgagtttattatca
ccaaacacaaaaatgcatacgacgcgtgggtgtgaactgcgcacagAATCCACCAGCTCGTG
AGAGACCCGAGGACAGGCCCTTCGACTACCGTGATTACAGCCGCTCGTCCGATGAT
GAACCGAGCAAGAGCACCGTCGCCGCCCTCCGGAGCAGGGGGCTTCGACTACGACAACACTACAGC
GGGGCCGACGAACGTCGTGGTGCACCGATGAATAACAAGGCGCCGAGCAGCAGCCTCGTGG
AGCGGGGCGTACATGGCTAGGGCGGAAGGCGGAGACGACGACGGTGTCTTCACGAGGAG
GCGGTGCGCGTCGGCAGGAGGCTCCATTCCACTTCCCGCCGGGACTCCGCCGCTCTGGT
TTCCCTGCCGCCAGGTGCGCCACTCCGTCCCGTTACGACGGCCGCGCTGCCGGCATCCTC
GCGACGTTGGCATCGCGTCCGACTCCACCCACGGTGCCAGCATGGAGGCGACGCTGCGCGCC
TGCGAGTCGCCACCATCGCCGGGAGTCCAAGTTCTGGCGACTTCGCTGGAGGGCCCTGGTG
GAGCGGCCATGGGAGTGCTGGGACCCGGGACATCAGGCCGGTGACGTCGACGCTGCCCGC
GCCGGCGCCCGCTGACGACGTACACCGTCGCGCCGTGACGCCGGTGAGGGGGGCGCTGTC
TTCGTGGCGTGCACGACGAGGCCTACCCGTACACCGTGTACCGGTGCCACACCACCGGCC
TCCAGGGCGTACACGGTGGACATGGAGGGCGCCGGCGACGCCGGTGACCATCGCCGCC
GTGTGCCACACCGACACGTCCTGTGGAACCCGGAGCACGTCTCCTTCAAGCTCCCTGGCACC
AAGCCCGGGCACGCCGGTGTGCCACCTCATGCCGTACGGGCACATAATCTGGGCCAAGAAC
GTGAAGCGCTGCCGGCGTGAGCGGCCTTGCAGCTCTGTGGTGTGCCGAACATAGATCGAT
GTACTACTACTATCTGTTCTACCTACGTCTTGTTCATACCACAGATGGTCACCCA
AGAGCAAGCGTTGTAATAAAAGAACAGCTTTGCAGAAGCTGGTGTATT

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Fig. 16. *taRAFTIN1b* promoter sequence (2095 bps).

TTGTTGAGTGCCACACTATTCACTACACCATAATGCACATTATGCTTGGATTGTCTTGTACT
 TGACTCATGTGTTAGACACTTCATTTATTTGGTGTGAATGACTCCTATGCTTACCATAGACCTTCATTGAGCGCTTGTGCATGTTGTTACACCTTGAGGTAGATGTTGTTCTCTTGTCAAATATATAGCATCTACCTCCCATTGCATGCTTGTACACCATATGCTAGGCTTGATGATGACACTTGTGGGTGACTCACCTTGAATGATTGGTTTGCAATTACGCTAACACACATTATTTTCCAAGTGTGTTGTCCTGCTCCTTGAAGGAACCACATGACGGTGCACATTGGAGAGTGCCTATTCGAGCTTCAAGATGATGAGTGTGTTGATCGTCCACTTCTACATGGTGACGCCGTCTTCCCATTGGTGAATTGGTTTGATCCGAGGTGGATCTTCCCAAGTGGGAGGGGATGATGCGGAGCATACTACGGACATCACCAGTCTAGAGTTCAATTACGCAAGTGACACCTTACACATCTACTACATAAAAGGTGAATCATCTCCTTACACGTGCTACTTGATCCCTTGAGGATGGTATACTACTTGACACTTCTCACGTGTGCATGCATAGGCATTGTCGGAGCACCATGAAACGATGAGGAGGGAGTGCAGACAAAGTGTACAACATACACCATCCGGAGGGAAAGCATGGAAGAGAAGGAAGAAGACATGGACAAGCTCTGGAAAGCCCGAACCTCTGGCCTCCTGGCCAAATCTCTGGATAGCCGGACCTCGACCCGAACCTCCGGCGCTGGACCTTCCGGCCATCCCTGGAACTCCCGGCCCTGCGCTATCCCTCGACAGACTCGGGCCGAAGCCGATGTACCCCTTCTGGCCCTCACTTATCCCTTCGTTGCTATCACTATATATACTCATCCTCCTCCATTCTAGGGTTAGCATTTGATAGCTCATTCGATGTGAGATTGCTCCTTACCCCCATCTCCTTGTGAGAGAGTGGAGATTGATGCACTCCATTGGAGTCCAAGGTCTCTTGGAGAAGATCCCATTGGGAATCAAGACCCCATCATGGGAAGATCCTCTAGGATTCAAGACCTCAACTCCTTAAGGATTGGGATGAACTAGTTACCTCTGTATCTCTGTGTTGGATTAAACCTTGTATCCCTCATGTTGATGTGGATTAGCATATGTGTGATTGGATCTTGTCTATTGGAGTGTGTTCCCTCTTTGTTTCTGTGTTCATCGTTTCTCGGGAGATCCCCCTCAATTCTGAAAGATCGGTCCCTAGGGTTCTACCCCTACATTAGCTCAGGTTCCCTACACATCTCGTTGTGAGCTGTTGCGCTTCTACGGCTGGGAGCTAAGCACATCTCATTCCCACCAAACGGGGTTCTCACATTGTAAACTTCATCGTATTTGCGAACTGCTTCTGGGACAGCCACTCACTTGAGTTGCTCCGATACTCTTCCGGGTCTCGTTCAGACCAACGGGGACACCGTCTGCAACCTGGAGGAGCCATTCTGCGACACACCAAAATTTCGCACGGACCCCCCGAAGATCCGCAAGAAAAAAAGCTGCAACGGCGTGGACGGCGAGCACCGCACCGCACAGCAACAGCGACGCGTACGCCCACGATAATATTCTCCGGTGCCTGACGTACCATGCGATCGCACAGCTCACCGATGTCACTCGCCACGATAATATTCTCCGGTGCCTGACGTACCATGCGATCGCACAGCTCACCGAGAGCTTCTGTTGGTGTGCCGTCAATGAAACACCTTCCCGTCAAGCCGACGACGCCATTAAAGTACCTCGCCTGATCGCATTACTACCTCCAAAGTACTACAAACCTCTCGACCTCTCACCTAGCGCACATCCATG

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Fig. 17. *taRAFTIN1d* predicted cDNA sequence (246 bps).

ATGGCGCGCTTCCTCGCCCTCCTCGCTGCCACCCCTGGTCGGGTTCAAGGCTGGAGGGCAG
CTGGGCCACGCAGCGCCGGCGACGGCGGAGGTGTTCTGGCGCCGTGCTGCCGACTGCCA
TTGCCCGACGCCGTTCTCCGCCTCTCAAACAAACCTGCAGCAGGTGTTGAACTGCACACAGAA
GCCACCAGCTCGTAAGAGACCCCGAGGAACAGGGCCCCCTCGACTACCGTGATTAC

Fig. 18. *taRAFTIN1d* partial genomic sequence (441 bps). Introns are shown in lower case letters.

ATGGCGCGCTTCCTCGCCCTCCTCGCTGCCACCCCTGGTCGCGgtaatggccgaagaagcc
actgagcaacgcctgcacatcttctttatttggcaaactggctaacggccaatactgcccgt
tgcgttacgtctcagGTTCAGGCTGGAGGGCAGCTGGGCCACGCAGCGCCGGCGACGGCGGAG
GTGTTCTGGCGCCGCGCTGCTGCCGCACTCGCCATTGCCGACGCCGTTCTCCGCCTCCTCAAA
CAACCTGCAGCAGgtctgtcttgcacatgttcctcgccctccgttaactgtttttctctc
gagtttgcattgtatcaccaaaacacaaaaatgcacatgcacgcgtacgcgttagGTGTTGAACGCAC
ACAGAACGCCACCAGCTTCGTAAGAGACCCCCGAGGACAGGGCCCCCTTCGACTACCGTGATTAC

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Fig. 19. *osRAFTIN1* cDNA (1301 bps, ORF from nt 63 to nt1301). Start and stop codons are underlined.

GTGCGAGTCGTCTCCGGGAGAAAATCGGCTGCGCCCCGTCTCTCTCTCTCGAACGCTTCCA
TGGCGCGCTTCCTCCTCCTCGTCGCCGTGCCGCTGCCGCCGTGCTTCTGCTGGCG
ACCGGGCGCGTCGACGGCCGAGGTGTTCTGGCGGCCGTGCTGCCGAATCCCCGTTGCCGG
ACGCCTTCCTCCGCCCTCCGCCCTGACACCAGCTTCGTCGTCGGCAAAGCGGAGGCGGCCG
GTGGCGCGGCCGCGGACCGGATTCCCCCTCGATTACACTGACTACAGGGGATCTGATTCTCCGA
CGACGGCGAGTGGTTGGACCTCGCCGGTGACTTCGGCGAGCCGGCGCCCTTCGGCTACGACT
ACAGTGCACAGGGCGAAGGCGGCGGCCGCCGCCGCCGCCGGAGAGCAGGTTCTG
CCGTCACGCGGGCTCAACTACGACAAATACGTCGGCGAGGAAGCTCCGCGGGCAGCA
GCACCGCCGGCGGAGAGAATGATGACGAGCCTTCGGGTACGACTACAAGGCGCCAGCAGCG
GCAGCGGCACCGCGGGCGTCGACGACGGCGAGGGCGTCGGCACGGCGCCACGACGACGGTGT
TCTTCCACGAGGAGGCGGTGCGCGTCCGCGAGAGGCTCCGTTACTTCCCAGGGCGACGA
CGTCGGCGCTGGGCTTCCTGCCGCCGCGTCGGACTCCATCCGTTCACGGCGGCCGCG
TGCCGGCCGTCTCGCGTGTTCGGCGTCGCGCCGGACACCGCCGAGGGCGGCCATGAGGG
AGACGCTGCGCACGTGCGAGTGGCCGACCCCTCGCCGGAGTCAAAGTTCTGCGCCACGTCGC
TGGAGGCCCTGGTGGAGGGGCCATGGCGCGCTCGGGACACGCGACATCGCCGCGCTGGCGT
CGACGCTGCCCGCGGCCGCCGCGCTGCAGGCGTACGCCGTCGCGCCGTGCTCCCCGTG
AGGGCGCCGGCTTCGTGGCGTGCACGACCAGGGCGTACCCGTACACCGTGTACCGCTGCCACA
CCACCGGCCGGCCAGAGCTTACATGGTGGAGATGGAAGGCGACGGCGGGCGATGGCGCG
AGGCGGTGACCGTGGCCACCGTGTGCCACACCAACACGTCGCGGTGGAACCCGGAGCACGTCT
CGTCAAGCTCCTCGGCACCAAGCCGGCGGCTGCCGGTGTGCCACCTCATGCCGTACGGGC
ACATCGTCTGGGCAAGAACGTGAAGAGCTGACGGCGTAG

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Fig. 20. *OsRAFTINI* genomic sequence (1479 bps, two introns included). Introns are shown in lower case letters.

GTGCGAGTCGTCTCCGGCGAGAAATCGGCTGCGCCCCGTCTCTCTCTCTCGAACGCTTCCA
TGGCGCGCTTCCTCCTCCTCGTCGCCGTGCCGCTGCCGCCGTGCTTCGgtacact
catgatgccgtactcagctgagccatgcaccgttgcacccgtataactaacgatcgctcgatc
gaccgacgtgtgtgtttttcagcagCTGGGCGACGCCGCGCCGTGACGCCGAGGTGTCT
GGCGCGCCGTGCTGCCGGAATCCCCGTTGCCGGACGCCCTCCTCCGCCCTCCGCCCTGgtc
ggtgtccttccttccttcgcgcgcgcgcgcgcattactctcctcgaggtttgatgg
tttgtggacgttgca^gACACCAAGCTTCGTCGGCAAAGCGGAGGCCGGTGGCGCG
CGGACCGGATTCCCCCTTCGATTACACTGACTACAGGGGATCTGATTCTCCGACGACGGCGAGT
GGTTTGGACCTCGCCGGTGACTTCGGCAGGCCGCGCCCTTCGGCTACGACTACAGTGCACAG
GGCGAAGGCCGGCGGCCGGCGCCGCGCCGCCGCCGGAGAGCAGGTTCTGCCGTCACGCCG
GGCTTCAACTACGACAAATACGTCGGCGAGGAAGCTCCGCGGGCAGCAGCACGCCGGC
GGAGAGAATGATGACGAGCCTTCGGGTACGACTACAAGGCGCCAGCAGCGGAGCGG
GCAGCGTCGACGACGGCGCGAGCGTCCGGCACGGCGCCACGACGACGGTGTCTCCACGAG
GAGGCCGTGCGTCCGGCGAGAGGCTCCGTTCTACTTCCGCCGGCAGCAGCTGGCGCTG
GGCTTCCCTGCCGCCGCCGCGTCGGACTCCATCCGTTCACGGCGGCCGCGCTGCCGGCGTC
CTCGCGCTGTTCCGGCGTCGCCGCCGGACACCGCCGAGGCCGGCGCATGAGGGAGACGCTGCGC
ACGTGCGAGTGGCCGACCCCTGCCGGCGAGTCCAAGTTCTGCCACGTCGCTGGAGGCCCTG
GTGGAGGGGCCATGGCGGGCGTCGGGACACGCGACATGCCGCCGCTGGCGTCAGCTGCC
CGCGGCCGGCGCCGCGTCAGCGTACGCCGCGCCGTGCTCCCCGTGAGGGCGCCGGC
TTCGTGGCGTGCCACGACCAGCGTACCCGTACACCGTGTACCGCTGCCACACCACGGCCCG
GCCAGAGCTTACATGGTGGAGATGGAAGGCGACGGCGGCCGATGGCGGCGAGCGGTGACC
GTGGCCACCGTGTGCCACACCAACACGTCGCCGGTGGAACCCGGAGCACGTCTCGTTCAAGCTC
CTCGGCACCAAGGCCGGCGTCGCCGGTGTGCCACCTCATGCCGTACGGCACATCGTCTGG
GCCAAGAACGTGAAGAGCTGACGGCGTAG

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Fig. 21. *osRAFTIN1* promoter sequence (1461 bps).

CGAAGGCAAACCTGGTAAGGATTCCCATACAGAACATCAATTATAAGTCTAAAACGAACA
CTATGTTATGAGAAACACCTCACATCCGTCCATAACCGTGGCATGACTATTAAAAAGTTA
ACTAAACTCTACAAAAGTTGCACGCTTACCCACACGTATGAACGTTCACATTACCGAATA
CATGTGGATCGGACATGGCCGACAAAGGAGAGTTCAATACAAGGCTTTCCATAACCAATCCA
TAAATATCCTATGTCCCACGGTGGGAACTCTCCACCAAACATCAAGCCAGGATCAGGT
CCTCATCTACCCATGCCACTCCATGGACTCCGACACATCCCACTGCAGGAGATTGCCATA
TACGCCACCATACCAGTGCTCCTCAACCGCTAACATGTTGGACACCAAATTCTATATACTTAT
ATAGTTCATCTCCACTAAGTGTAGTTAATTACATTCTCTCTCTCATTAAGCCACATCAC
CTCAATTATTTAGCCTTAGATGATAGATCTATGGCCAATTGTCTTTCTTCTTCT
CTTAAAAACATGCAATCTTAAACTTTAGGCTAAAATTGTATCAAATTGTTTAGTTG
TACATATTATGCAACTTAATTTCGCCGCAACCGGGAGGGTATTCATCTAGTATTATTTA
AGAGCTATACACACTGCTATAGGGGAAAAAAAGATAGGTTGGCCCCCTGGTCAGTCCTGTT
GCACGGCTATATGTTGAAGGGAAAAGCCAGTACGTTTGTAGGTTGTTTTAGAATT
GCTAAAAAGTTGGCATGTTTTAGGTTAAAGCCTTAAATATAAGTTACATTGTAACATAC
AGTGTAACTTCCGCTGTAACATATTGTAATCTCTATATAAGTTAGATATAAAATTACATATAT
ATTATTTAATACCTATTATAAGTTAGTATATTAGTTATAATGGAATTAAATTATAATTAT
AGTATAGTTAGATTGAAAGTTTTCTTTAAGAAATTTCGCAACAGTTATTAGATATAGTC
CCTAAACGAAAATGTCAGGTGGATGCATGATTCACTGTCAGTGTGACGCTCGGGCGGATCACGGCTGCG
TCACGAAAATTCCCCCATGCAACCCCGGTCCGGCCGTCCTCGTGCCAACAGGCAACAGCGC
GGCGCCGGCGAACGTACGCCAAGATTATATTCCCCCTCTCGCGCTCGCGCGCCGCGACG
TCGTCGGAGCCAACATTATTTCGTTCTGTCACCGTCGCCGTTGATCTCAAGCGAGATT
TGAGGTTGCCACGACGACGCCCTGCTATAAATACCAAGGTGGTGGTCACCGCCGGCGCGT
CGATCGATCCGTCGCAGTCGTCTCCGGCGAGAAATCGGCTGCGCCCCGTCTCTCTCG
AACGCTTCCATG

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Fig. 22. Predicted protein sequences

taRAFTIN1a (389 residues)

MARFLVALLATTAVQAGGQLGHAAPATAEVFWRAVLPHSPLPDAVLRLLKQPAAGVELLTEATSFVR
DAEDRPPFDYRDYSRSPPDDEPSKSTGAASGARDFDYDDYSGGDKLRAASGARDFDYDDYSGADKLRG
ATDEYKAPSSSLAGNGASMARGGKAETTTVFFHEEAVRVGKRLPFRFPPATPAALGFLPRQVADSPVFT
TAALPGVLATFGVASDSATVASMEATLRACESPTIAGESKFCATSLEALVERAMEVLGTRDIRPVTSTL
PRAGAPLQTYTVRSVRPVEGGPVFVACHDEAYPYTVRCHTTGPSRAYVDMEGARGGDAVTIATVCHT
DTSLWNPEHVSFKLLGTPKGTPVCHLMPYGHIIWAKNVNRSWA

taRAFTIN1b (362 residues)

MARFLVALLAATLVAVQAGGQLGHAAPATGEVFWRAVLPHSPLPDAVLRLLKQPAAEESTSFVRDPEDRP
PFDYRDYSRSSDDEPSKSTVAASGAGGFDYDNYSGADERRGATDEYKAPSSSLAGSGAYMARGGKAET
TTVFFHEEAVRVGRRLPFHFPPATPAALGFLPRQVADSPVFTTAALPGILATFGIASDSTVPSMEATL
RACESPTIAGESKFCATSLEALVERAMGVLGTRDIRPVTSTLPRAGAPLQTYTVVAVQPVEGGPVFVAC
HDEAYPYTVRCHTTGPSRAYVDMEGARGADAVTIAAVCHTDTSLWNPEHVSFKLLGTPKGTPVCHL
MPYGHIIWAKNVKRSPA

taRAFTIN1d (partial sequence, 82 residues)

MARFLVALLAATLVAVQAGGQLGHAAPATAEVFWRAVLPHSPLPDAVLRLLKQPAAGVELHTEATSFVR
DPEDRPPFDYRDY

osRAFTIN1 (412 residues)

MARFLLLLVAVAAAAAVSLGDAAPSTAEVFWRAVLPESPPLPDAFLRLLRPDTSFVVGKAEAGGAART
GFPFDYTDYRGSDSPTTASGLDLAGDFGEPAFPFYDYSAQGEGGGGGAAAAAGEQVLAVDAGFNYDKYV
GARKLRGGSSTAGGENDDEPFYDYSQKAPSSGSGTAASTTARGVGTGATTTVFFHEEAVRVGERLPFYFP
AATTSALGFLPRRVADSIPFTAALPAVLALFGVAPDTAEAGMRETLRTCEWPTLAGESKFCATSLEA
LVEGAMAALGTRDIAALASTLPRGGAPLQAYAVRAVLVPEGAGFVACHDQAYPYTVRCHTTGPARAYM
VEMEGDGGGDGGEAVTVATVCHTNTSRWNPEHVSFKLLGTPKGSPVCHLMPYGHIVWAKNVKSSTA

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